

SEQUENCE LISTING

<110> Andrew, David P.
Zabel, Brian A.
Ponath, Paul D.

<120> ANTI-GPR-9-6 ANTIBODIES AND METHODS OF
IDENTIFYING AGENTS WHICH MODULATE GPR-9-6 FUNCTION

<130> LKS98-16

<140> 09/266,464

<141> 1999-03-11

<160> 7

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<211> 2577

<212> DNA

<213> Homo sapiens

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Asn Phe Asn Phe Thr Asp Phe Tyr Cys Glu Lys Asn Asn Val Arg Gln	
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Phe Ala Ser His Phe Leu Pro Pro Leu Tyr Trp Leu Val Phe Ile Val	
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Gly Ala Leu Gly Asn Ser Leu Val Ile Leu Val Tyr Trp Tyr Cys Thr	
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Arg Val Lys Thr Met Thr Asp Met Phe Leu Leu Asn Leu Ala Ile Ala	
70 75 80	

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gac Asp	cag Gln	tgg Trp 100	aag Lys	ttc Phe	cag Gln	acc Thr	ttc Phe 105	atg Met	tgc Cys	aag Lys	gtg Val	gtc Val 110	aac Asn	agc Ser	atg Met	396
tac Tyr	aag Lys 115	atg Met	aac Asn	ttc Phe	tac Tyr	agc Ser 120	tgt Cys	gtg Val	ttg Leu	ctg Leu	atc Ile 125	atg Met	tgc Cys	atc Ile	agc Ser	444
gtg Val 130	gac Asp	agg Arg	tac Tyr	att Ile	gcc Ala 135	att Ile	gcc Ala	cag Gln	gcc Ala	atg Met 140	aga Arg	gca Ala	cat His	act Thr	tgg Trp 145	492
agg Arg	gag Glu	aaa Lys	agg Arg	ctt Leu 150	ttg Leu	tac Tyr	agc Ser	aaa Lys	atg Met 155	gtt Val	tgc Cys	ttt Phe	acc Thr 160	atc Ile	tgg Trp	540
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325 330 335

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Glu Gly Ser Leu Lys Leu Ser Met Leu Leu Glu Thr Thr Ser Gly
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Val Gly Ala Leu Gly Asn Ser Leu Val Ile Leu Val Tyr Trp Tyr Cys
50 55 60

Thr Arg Val Lys Thr Met Thr Asp Met Phe Leu Leu Asn Leu Ala Ile
 65 70 75 80
 Ala Asp Leu Leu Phe Leu Val Thr Leu Pro Phe Trp Ala Ile Ala Ala
 85 90 95
 Ala Asp Gln Trp Lys Phe Gln Thr Phe Met Cys Lys Val Val Asn Ser
 100 105 110
 Met Tyr Lys Met Asn Phe Tyr Ser Cys Val Leu Leu Ile Met Cys Ile
 115 120 125
 Ser Val Asp Arg Tyr Ile Ala Ile Ala Gln Ala Met Arg Ala His Thr
 130 135 140
 Trp Arg Glu Lys Arg Leu Leu Tyr Ser Lys Met Val Cys Phe Thr Ile
 145 150 155 160
 Trp Val Leu Ala Ala Ala Leu Cys Ile Pro Glu Ile Leu Tyr Ser Gln
 165 170 175
 Ile Lys Glu Glu Ser Gly Ile Ala Ile Cys Thr Met Val Tyr Pro Ser
 180 185 190
 Asp Glu Ser Thr Lys Leu Lys Ser Ala Val Leu Thr Leu Lys Val Ile
 195 200 205
 Leu Gly Phe Phe Leu Pro Phe Val Val Met Ala Cys Cys Tyr Thr Ile
 210 215 220
 Ile Ile His Thr Leu Ile Gln Ala Lys Lys Ser Ser Lys His Lys Ala
 225 230 235 240
 Leu Lys Val Thr Ile Thr Val Leu Thr Val Phe Val Leu Ser Gln Phe
 245 250 255
 Pro Tyr Asn Cys Ile Leu Leu Val Gln Thr Ile Asp Ala Tyr Ala Met
 260 265 270
 Phe Ile Ser Asn Cys Ala Val Ser Thr Asn Ile Asp Ile Cys Phe Gln
 275 280 285
 Val Thr Gln Thr Ile Ala Phe Phe His Ser Cys Leu Asn Pro Val Leu
 290 295 300
 Tyr Val Phe Val Gly Glu Arg Phe Arg Arg Asp Leu Val Lys Thr Leu
 305 310 315 320
 Lys Asn Leu Gly Cys Ile Ser Gln Ala Gln Trp Val Ser Phe Thr Arg
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 Gly Ala Leu Ser Leu
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<212> DNA

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